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TECH CENTER 1600/2800

**ENTERED** OIPE**RAW SEQUENCE LISTING**

PATENT APPLICATION: US/09/936,145

DATE: 03/15/2002

TIME: 14:44:04

Input Set : A:\011309.txt

Output Set: N:\CRF3\03152002\I936145.raw

#9

```

3 <110> APPLICANT: Inoue, Yasushi
4      Fushimi, Naoya
5      Mizubuchi, Hiroyuki
6      Yamamoto, Yoshie
7      Ohshima, Yoshie
8      Yasutake, Nozomu
9      Miyoshi, Shinsuke
11 <120> TITLE OF INVENTION: Promoters
13 <130> FILE REFERENCE: 3274-011309
15 <140> CURRENT APPLICATION NUMBER: 09/936,145
16 <141> CURRENT FILING DATE: 2001-09-07
18 <150> PRIOR APPLICATION NUMBER: PCT/JP00/01415
19 <151> PRIOR FILING DATE: 2000-03-08
21 <150> PRIOR APPLICATION NUMBER: US11/060904
22 <151> PRIOR FILING DATE: 1999-03-08
24 <150> PRIOR APPLICATION NUMBER: US11/286034
25 <151> PRIOR FILING DATE: 1999-10-06
27 <160> NUMBER OF SEQ ID NOS: 22
29 <170> SOFTWARE: Microsoft Word 97 SR-2
31 <210> SEQ ID NO: 1
32 <211> LENGTH: 249
33 <212> TYPE: DNA
34 <213> ORGANISM: Bacillus amyloliquefaciens
36 <400> SEQUENCE: 1
37 gccgcgacac tacgaaaaga ctggctgaaa acattgagcc tttgatgact gatgatttgg      60
38 ctgaagaagt ggatcgattg tttgagaaaa gaagaagacc ataaaaatac cttgtctgtc      120
39 atcagacagg gtatttttta tgctgtccag actgtccgct gtgtaaaaaa taggaataaa      180
40 ggggggttgt tattatttta ctgatatgta aaatataatt tgtataagaa aatgagaggg      240
41 agaggatcc                                     249
43 <210> SEQ ID NO: 2
44 <211> LENGTH: 270
45 <212> TYPE: DNA
46 <213> ORGANISM: Bacillus amyloliquefaciens
48 <400> SEQUENCE: 2
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50 ctgaagaagt ggatcgattg tttgagaaaa gaagaagacc ataaaaatac cttgtctgtc      120
51 atcagacagg gtatttttta tgctgtccag actgtccgct gtgtaaaaaa taggaataaa      180
52 ggggggttgt tattatttta ctgatatgta aaatataatt tgtataagaa aatgagaggg      240
53 agaggatcc ccgggtaccga gctcgaattc                                     270
55 <210> SEQ ID NO: 3
56 <211> LENGTH: 29
57 <212> TYPE: DNA
58 <213> ORGANISM: Artificial Sequence

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60 <220> FEATURE:
61 <223> OTHER INFORMATION: Description of Artificial Sequence: Primer
63 <400> SEQUENCE: 3
64 cgctctagag ccccgcacat acgaaaaga 29
66 <210> SEQ ID NO: 4
67 <211> LENGTH: 35
68 <212> TYPE: DNA
69 <213> ORGANISM: Artificial Sequence
71 <220> FEATURE:
72 <223> OTHER INFORMATION: Description of Artificial Sequence: Example of a primer
73     for introducing a restriction site
75 <400> SEQUENCE: 4
76 cgcgaattcg gatcctctcc ctctcatttt cttat 35
78 <210> SEQ ID NO: 5
79 <211> LENGTH: 50
80 <212> TYPE: DNA
81 <213> ORGANISM: Artificial Sequence
83 <220> FEATURE:
84 <223> OTHER INFORMATION: Description of Artificial Sequence: Example of a primer
85     for introducing a restriction site
87 <400> SEQUENCE: 5
88 cgcgaattcg agctcggtac ccggggatcc tctccctctc attttcttat 50
90 <210> SEQ ID NO: 6
91 <211> LENGTH: 29
92 <212> TYPE: DNA
93 <213> ORGANISM: Artificial Sequence
95 <220> FEATURE:
96 <223> OTHER INFORMATION: Description of Artificial Sequence: Primer
98 <400> SEQUENCE: 6
99 cgcggatcca tgtattacaa caggttggt 29
101 <210> SEQ ID NO: 7
102 <211> LENGTH: 29
103 <212> TYPE: DNA
104 <213> ORGANISM: Artificial Sequence
106 <220> FEATURE:
107 <223> OTHER INFORMATION: Description of Artificial Sequence: Primer
109 <400> SEQUENCE: 7
110 cgcgaattct cacacatact ccttcgtat 29
112 <210> SEQ ID NO: 8
113 <211> LENGTH: 29
114 <212> TYPE: DNA
115 <213> ORGANISM: Artificial Sequence
117 <220> FEATURE:
118 <223> OTHER INFORMATION: Description of Artificial Sequence: Primer
120 <400> SEQUENCE: 8
121 cgcggatcca tgtcttggtc aattagctc 29
123 <210> SEQ ID NO: 9
124 <211> LENGTH: 29
125 <212> TYPE: DNA

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126 <213> ORGANISM: Artificial Sequence
128 <220> FEATURE:
129 <223> OTHER INFORMATION: Description of Artificial Sequence: Primer
131 <400> SEQUENCE: 9
132 aaagaattct taatcaacac gcccgttat 29
134 <210> SEQ ID NO: 10
135 <211> LENGTH: 26
136 <212> TYPE: DNA
137 <213> ORGANISM: Artificial Sequence
139 <220> FEATURE:
140 <223> OTHER INFORMATION: Description of Artificial Sequence: Primer
142 <400> SEQUENCE: 10
143 gtttcctctc cctctcattt tcttat 26
145 <210> SEQ ID NO: 11
146 <211> LENGTH: 20
147 <212> TYPE: DNA
148 <213> ORGANISM: Artificial Sequence
150 <220> FEATURE:
151 <223> OTHER INFORMATION: Description of Artificial Sequence: Primer
153 <400> SEQUENCE: 11
154 atgtattaca acaggttggt 20
156 <210> SEQ ID NO: 12
157 <211> LENGTH: 20
158 <212> TYPE: DNA
159 <213> ORGANISM: Artificial Sequence
161 <220> FEATURE:
162 <223> OTHER INFORMATION: Description of Artificial Sequence: Primer
164 <400> SEQUENCE: 12
165 atgtcttggt caattagctc 20
167 <210> SEQ ID NO: 13
168 <211> LENGTH: 29
169 <212> TYPE: DNA
170 <213> ORGANISM: Artificial Sequence
172 <220> FEATURE:
173 <223> OTHER INFORMATION: Description of Artificial Sequence: Primer
175 <400> SEQUENCE: 13
176 cgcgaattca tgtattacaa caggttggt 29
178 <210> SEQ ID NO: 14
179 <211> LENGTH: 29
180 <212> TYPE: DNA
181 <213> ORGANISM: Artificial Sequence
183 <220> FEATURE:
184 <223> OTHER INFORMATION: Description of Artificial Sequence: Primer
186 <400> SEQUENCE: 14
187 cgcgaattca tgtcttggtc aattagctc 29
189 <210> SEQ ID NO: 15
190 <211> LENGTH: 1581
191 <212> TYPE: DNA
192 <213> ORGANISM: Agrobacterium radiobacter M36

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194 <220> FEATURE:
195 <221> NAME/KEY: Promoter
196 <222> LOCATION: 314..316
198 <220> FEATURE:
199 <221> NAME/KEY: Terminator
200 <222> LOCATION: 1559..1561
202 <220> FEATURE:
203 <221> NAME/KEY: Gene
204 <222> LOCATION: 341..1558
205 <223> OTHER INFORMATION: MIase structural gene
207 <400> SEQUENCE: 15
208 gatctgcgtg cccatggcac cgtcgagaat gaggatgcgt tcgctggcag cctcgcgcag      60
209 cgccttgaaa atttcgcgc cgtcgcgctt tgccccttca gggccaaaca gatcgtcaaa      120
210 cacgggcaca ctctcatatt cgatttgcaa gatcgcaagt cgtcaagtca cataaagata      180
211 tgtttatgtc aatatactct caagggacag gcatggcttt gcgtcgttgc gtcacgttac      240
212 gaaatatcgc tgacagatga caggtttata cggcaaggat ataagccgaa gcagcaaacy      300
213 catggaggac gcaatgcccg aagacgatca caacagccgc aactggaata cctgcacctg      360
214 gcaccgccag tggctggtga aacaggccga gggacttttc gacttcttcc agtatcgcgc      420
215 cctcaatccc gccggcggtt tcttcgatct cgacgccaa ggcgcgcgcgc tgcaggcaaa      480
216 cgatcccgct gcgggcatcc atgcctctgc gcgcatggtg cattgcttct ccatcggcca      540
217 cctgctcggc cggcggggct gcggcgatat cgtcgaccac ggcattgacct atctctggaa      600
218 caaacaccgc gatggcgaac atggcggtta tttctggcag gtgaacgatg ccggcccagt      660
219 ggacgccacc aagcagggtt atggccacgc ctctcgtgct ctggccgcct ctccgcgcaa      720
220 gaccgtcggc caccgcgtgg ccgaccggat gctggctgat attaccgaag tgctggaaag      780
221 tcgtttctgg gaagaaaaac atggcgccat cgccgaggaa ttcaatcgcg actggtcgcc      840
222 catcgacaat tatcgcggcc agaactccaa tatgcacctc accgaagcgc tgatggccgc      900
223 ctatgaggtg accggcgaca ataactatct cagcaaggcc gaacgcatcg ccgatctcgt      960
224 catccgtcgc cgcgcgcggc agctggattt ccgcgtgccc gagcatttcg acgacaactg     1020
225 gacgtctggc aaggactatc gcggcaacga aatgttccgc ccctccggct ccacccccgg     1080
226 cactggctg gaatggcgcc gtctcatcct gcaattgtgg atactgggcg aacgcgcgca     1140
227 cgactggatg ccggtcgcgg ccaaatccct ctctcgtcag tccatggcgc tgggctggga     1200
228 ccgggaaaag ggcggcttct tttatacgtt ggactggaat gacaatcccg acaagcgggc     1260
229 aaagctctgg tggcccatgt ccgaggcggc ggggtcggcc catttctca acgagaacct     1320
230 gccggcggtt ggcttctacg aagacagcta tcgtcgcata tggagcacca tcgccaacaa     1380
231 ctccatcgac catgccaatg gcggctggca tgaggaaact acggaagatc tggttccgcg     1440
232 ccacacgcta ttcccaggca agggcgatat ctaccatgcg ctccaggcct gcctcatccc     1500
233 gcttttcccg gcgacgggca gcctgacgaa ggtgatcaag gaaagcggcg gggattatta     1560
234 aggcgtctg cgccaatag c                                     1581
236 <210> SEQ ID NO: 16
237 <211> LENGTH: 39
238 <212> TYPE: DNA
239 <213> ORGANISM: Artificial Sequence
241 <220> FEATURE:
242 <223> OTHER INFORMATION: Description of Artificial Sequence: Primer
244 <400> SEQUENCE: 16
245 gcatctcgag catatgcgga tcctctccct ctcatcttc      39
247 <210> SEQ ID NO: 17
248 <211> LENGTH: 31
249 <212> TYPE: DNA

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250 <213> ORGANISM: Artificial Sequence
252 <220> FEATURE:
253 <223> OTHER INFORMATION: Description of Artificial Sequence: Primer
255 <400> SEQUENCE: 17
256 gcatctcgag ggtaataaaa aaacacctcc a 31
258 <210> SEQ ID NO: 18
259 <211> LENGTH: 30
260 <212> TYPE: DNA
261 <213> ORGANISM: Artificial Sequence
263 <220> FEATURE:
264 <223> OTHER INFORMATION: Description of Artificial Sequence: Primer
266 <400> SEQUENCE: 18
267 gcatgaattc aaagcagcga tcccgatgaa 30
269 <210> SEQ ID NO: 19
270 <211> LENGTH: 283
271 <212> TYPE: DNA
272 <213> ORGANISM: Bacillus amyloliquefaciens
274 <400> SEQUENCE: 19
275 ctcgagggta ataaaaaac acctccaagc tgagtgcggg taccagcttg gaggtgcgtt 60
276 ttttttttca gccgtatgac aaggctcgga tcaggtgtga caaatacggg atgctggctg 120
277 tcataggtga caaatccggg ttttgcgcg tttggctttt tcacatgtct gatttttgta 180
278 taatcaacag gcaaggagcc ggaatctttc gccttgga aaataagcggc gatcgtagct 240
279 gcttccaata tggattgttc atcgggatcg ctgctttgaa ttc 283
281 <210> SEQ ID NO: 20
282 <211> LENGTH: 28
283 <212> TYPE: DNA
284 <213> ORGANISM: Artificial Sequence
286 <220> FEATURE:
287 <223> OTHER INFORMATION: Description of Artificial Sequence: Primer
289 <400> SEQUENCE: 20
290 gcatcatatg cccgaagacg atcacaac 28
292 <210> SEQ ID NO: 21
293 <211> LENGTH: 31
294 <212> TYPE: DNA
295 <213> ORGANISM: Artificial Sequence
297 <220> FEATURE:
298 <223> OTHER INFORMATION: Description of Artificial Sequence: Primer
300 <400> SEQUENCE: 21
301 gcatctcgag ttaataatcc ccgccgcttt c 31
303 <210> SEQ ID NO: 22
304 <211> LENGTH: 21
305 <212> TYPE: DNA
306 <213> ORGANISM: Artificial Sequence
308 <220> FEATURE:
309 <223> OTHER INFORMATION: Description of Artificial Sequence: Primer
311 <400> SEQUENCE: 22
312 atgcccgaag acgatcaca c 21

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VERIFICATION SUMMARY

PATENT APPLICATION: US/09/936,145

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Input Set : A:\011309.txt

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